



65446-87.txt
SEQUENCE LISTING

<110> Kane, Michael D.
Nagel, Aaron C.
Dombkowski, Alan A.

<120> COMPOSITIONS AND SYSTEMS FOR IDENTIFYING AND COMPARING EXPRESSED GENES
(mRNAs) IN EUKARYOTIC ORGANISMS

<130> 65446-87

<160> 5

<170> PatentIn version 3.1

<210> 1

<211> 61

<212> DNA

<213> Artificial Sequence

<220>

<223> This is an artificial sequence.

<220>

<221> misc_feature

<222> (1)..(50)

<223> The poly-T tail may have between 8 and 50 residues.

<220>

<221> misc_feature

<222> (51)..(51)

<223> v may be selected from a or c or g.

<220>

<221> misc_feature

<222> (52)..(62)

<223> The sequence may have between 3 and 10 residues selected from a o
r c or g or t.

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 n 61

<210> 2

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> This sequence is artificial.

<220>

<221> misc_feature

<222> (23)..(25)

<223> n may be selected from a or c or g or t.

<220>

<221> misc_feature

<222> (24)..(24)

<223> v may be selected from a or c or g.

<400> 2
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<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> This is an artificial sequence.

<400> 3

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24

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<212> DNA

<213> Artificial Sequence

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<223> This is an artificial sequence.

<400> 4

gtttaaaaaa aaaaaaaaaa aaaa

24

<210> 5

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<212> DNA

<213> Artificial Sequence

<220>

<223> This is an artificial sequence.

<400> 5

tttttttttt tttttttttt ggttt

25